## EXHIBIT B MARKED UP VERSION OF AMENDED CLAIMS

- 1. (Twice Amended) An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:
  - (a) the nucleotide sequence as set forth in SEQ ID NO: 1;
- (b) a nucleotide sequence encoding the polypeptide set forth in SEQ ID NO: 2;
- (c) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of (a) or (b); and
  - (e)—a nucleotide sequence complementary to any either of (a)—(e) or (b).
- 2. (Twice Amended) An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:
- (a) a nucleotide sequence encoding a polypeptide that is at least about 70, 75, 80, 85, 90, 95, 96, 97, 98, or 99 percent identical to the polypeptide set forth in SEQ ID NO: 2, wherein the encoded polypeptide has human E3 $\alpha$  ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (b) a nucleotide sequence encoding an allelic variant or splice variant of the nucleotide sequence as set forth in SEQ ID NO: 1, encoding a polypeptide that has human E3α ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (c) a nucleotide sequence of SEQ ID NO: 1; (a); or (b) encoding a polypeptide fragment of at least about 25 amino acid residues, wherein the polypeptide has an activity of the polypeptide set forth in SEQ ID NO: 2;
- (d) a nucleotide sequence of SEQ ID NO: 1, or (a) (c) comprising a fragment of at least about 16 nucleotides;
- (e) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a) (d); and
  - (f)—a nucleotide sequence complementary to any of (a)-(e) b).

- 3. (Twice Amended) An isolated nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of:
- (a) a nucleotide sequence encoding a polypeptide set forth in SEQ ID NO: 2 with at least one conservative amino acid substitution, wherein the polypeptide has an human E3α ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (b) a nucleotide sequence encoding a polypeptide set forth in SEQ ID NO:
   2 with at least one amino acid insertion, wherein the polypeptide has an human E3α
   ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (c) a nucleotide sequence encoding a polypeptide set forth in SEQ ID NO: 2 with at least one amino acid deletion, wherein the polypeptide has  $\frac{1}{2}$  human  $\frac{1}{2}$  ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (d) a nucleotide sequence encoding a polypeptide set forth in SEQ ID NO:
   2 which has a C- and/or N-terminal truncation, wherein the polypeptide has an human E3α ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (e) a nucleotide sequence encoding a polypeptide set forth in SEQ ID NO: 2 with at least one modification selected from the group consisting of amino acid substitutions, amino acid insertions, amino acid deletions, C-terminal truncation, and N-terminal truncation, wherein the polypeptide has an human E3α ligase activity of the polypeptide set forth in SEQ ID NO: 2;
- (f) a nucleotide sequence of (a)-(e) comprising a fragment of at least about 16 nucleotides;
- (g) a nucleotide sequence which hybridizes under moderately or highly stringent conditions to the complement of any of (a)-(f); and
  - (h)—a nucleotide sequence complementary to any of (a)-(e).
- 59. (Twice Amended) A diagnostic reagent comprising a detectably labeled polynucleotide encoding the amino acid sequence set out in SEQ ID NO: 2; or

a fragment, variant or homolog thereof including allelic variants and or spliced variants thereof with human E3 $\alpha$  ligase activity.

- 60. (Amended) The diagnostic reagent of claim 58 59, wherein said labeled polynucleotide is a first-strand cDNA.
- 61. (Amended) A method for determine determining the presence of huE3α nucleic acids in a biological sample comprising the steps of:
- (a) providing a biological sample suspected of containing huE3α nucleic acids;
- (b) contacting the biological sample with a diagnostic reagent according to claim 59 under conditions wherein the diagnostic reagent will hybridize with  $huE3\alpha$  nucleic acids contained in said biological sample;
- (c) detecting hybridization between huE3\alpha nucleic acid in the biological sample and the diagnostic reagent; and
- (d) comparing the level of hybridization between the biological sample and diagnostic reagent with the level of hybridization between a known concentration of  $huE3\alpha$  nucleic acid and the diagnostic reagent.
- 62. (Amended) A method for detecting the presence of huE3 $\alpha$  nucleic acids in a tissue or cellular sample comprising the steps of:
- (a) providing a tissue or cellular sample suspected of containing  $huE3\alpha$  nucleic acids;
- (b) contacting the tissue or cellular sample with a diagnostic reagent according to claim 59 under conditions wherein the diagnostic reagent will hybridize with huE3α nucleic acids;
- (c) detecting hybridization between huE3 $\alpha$  nucleic acid in the tissue or cellular sample and the diagnostic reagent; and

(d) comparing the level of hybridization between the tissue or cellular sample and diagnostic reagent with the level of hybridization between a known concentration of  $huE3\alpha$  nucleic acid and the diagnostic reagent.